SEQUENCE LISTING

<110> Kiyosue, Yuko Sasaki, Hiroyuki Tsukita, Shoichiro Eisai Co., Ltd.

<120> CULTURED XENOPUS LAEVIS CELL LINES EXPRESSING MUTANT ADENOMATOUS POLYPOSIS COLI GENE

<130> 082368-002400US
<150> PCT/JP03/10434
<151> 2003-08-19
<150> JP 2002-241487
<151> 2002-08-22
<160> 9
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 2829
<212> PRT
<213> Xenopus laevis
<400> 1

Met Ala Ala Ala Ser Tyr Asp Gln Leu Val Lys Gln Val Glu Ala Leu 10 5 Thr Met Glu Asn Thr Asn Leu Arg Gln Glu Leu Glu Asp Asn Ser Asn 25 20 His Leu Thr Lys Leu Glu Thr Glu Ala Thr Asn Met Lys Glu Val Leu 40 45 Lys Gln Leu Gln Gly Ser Ile Glu Asp Glu Ala Met Ala Ser Ser Gly 55 Pro Ile Asp Leu Leu Glu Arg Phe Lys Asp Leu Asn Leu Asp Ser Ser 70 75 Asn Ile Pro Ala Gly Lys Ala Arg Pro Lys Met Ser Met Arg Ser Tyr 85 90 Gly Ser Arg Glu Gly Ser Leu Ser Gly His Ser Gly Glu Cys Ser Pro 105 Val Pro Val Gly Ser Phe Gln Arg Arg Gly Leu Leu Asn Gly Ser Arg 120 125 Glu Ser Ala Gly Tyr Met Glu Glu Leu Glu Lys Glu Arg Leu Leu 135 Ile Ala Glu His Glu Lys Glu Glu Lys Glu Lys Arg Trp Tyr Ala 155 150 Gln Leu Gln Asn Leu Thr Lys Arg Ile Asp Ser Leu Pro Leu Thr Glu 170 Asn Phe Ser Met Gln Thr Asp Met Thr Arg Arg Gln Leu Glu Tyr Glu 185 Ala Arg Gln Ile Arg Ala Ala Met Glu Glu Gln Leu Gly Thr Cys Gln 200 205

Asp Met Glu Lys Arg Val Gln Thr Arg Val Gly Lys Ile His Gln Ile

Glu Glu Glu Ile Leu Arg Ile Arg Gln Leu Leu Gln Ser Gln Val Ala

Glu Ala Ala Glu Arg Thr Pro Gln Ser Lys His Asp Ala Gly Ser Arg

215

230

220

235

				245					250					255	
Asp	Ala	Glu	Lys 260	Leu	Pro	Asp	Gly	Gln 265	Gly	Thr	Ser	Glu	Ile 270	Thr	Ala
Ser	Gly	Asn 275	Val	Gly	Ser	Gly	Gln 280	Gly	Ser	Ser	Ser	Arg 285	Ala	Asp	His
Asp	Thr 290	Thr	Ser	Val	Met	Ser 295	Ser	Asn	Ser	Thr	Tyr 300	Ser	Val	Pro	Arg
Arg 305	Leu	Thr	Ser	His	Leu 310	Gly	Thr	Lys	Val	Glu 315	Met	Val	Tyr	Ser	Leu 320
Leu	Ser	Met	Leu	Gly 325	Thr	His	Asp	Lys	Asp 330	Asp	Met	Ser	Arg	Thr 335	Leu
			340				_	Ser 345	_				350		
	•	355					360	Leu				365			
	370			_		375		Gly			380		_		
385					390			Ile		395			_	_	400
				405				Leu	410					415	_
			420					Trp 425					430		
	_	435	_	_			440	Pro				445			
	450					455		Lys			460	_			
465					470			Gly		475					480
				485				Gly	490			_		495	
			500					Met 505					510		
		515					520	Leu				52 5			
	530					535		Ser			540	_			
545					550			Leu		555	_		_		560
				565				Gly	570					575	
			580					Ser 585					590		
		595					600	Cys				605			
	610					615		Phe			620				
625					630			Ile		635			_	_	640
				645				Ala	650			_		655	
			660					Gln 665					670		
		675					680	Asn			_	685		_	
	690					695		Gln			700				
705					710			Ile		715	_				720
ΑΙΑ	Met	GТĀ	ser	725	АТА	А1а	Leu	Arg	730	Leu	Met	Ala	Asn	Arg 735	Pro

Ala Lys Tyr Lys Asp Ala Asn Ile Met Ser Pro Gly Ser Ser Val Pro 745 Ser Leu His Val Arg Lys Gln Lys Ala Leu Glu Ala Glu Leu Asp Ala 760 Gln His Leu Ser Glu Thr Phe Asp Asn Ile Asp Asn Leu Ser Pro Lys 775 Thr Thr His Arg Asn Lys Gln Arg His Lys Gln Asn Leu Cys Ser Glu 795 790 Tyr Ala Leu Asp Ser Ser Arg His Asp Asp Ser Ile Cys Arg Ser Asp 805 810 Asn Phe Ser Ile Gly Asn Leu Thr Val Leu Ser Pro Tyr Ile Asn Thr 820 825 Thr Val Leu Pro Gly Ser Ser Ser Pro Arg Pro Thr Met Asp Gly Ser 840 Arg Pro Glu Lys Asp Arg Glu Arg Thr Ala Gly Leu Gly Asn Tyr His 855 860 Ser Thr Thr Glu Ser Ser Gly Asn Ser Ser Lys Arg Ile Gly Ile Gln 870 875 Leu Ser Thr Thr Ala Gln Ile Ser Lys Val Met Asp Glu Val Ser Asn 890 885 Ile His Leu Val Gln Glu Asn Arg Ser Ser Gly Ser Ala Ser Glu Met 905 His Cys Met Ser Asp Glu Arg Asn Ser Gln Arg Lys Pro Ser Ser Asn 920 925 His Pro Gln Ser Asn Pro Phe Thr Phe Thr Lys Ala Glu Ser Ser Thr 935 940 Arg Gly Cys Pro Val Ala Phe Met Lys Met Glu Tyr Lys Met Ala Ser 950 955 Asn Asp Ser Leu Asn Ser Val Ser Ser Thr Glu Gly Tyr Gly Lys Arg 970 965 Gly Gln Val Lys Pro Ser Val Glu Ser Tyr Ser Glu Asp Asp Glu Ser 985 980 990 Lys Phe Phe Ser Tyr Gly Gln Tyr Pro Ala Gly Leu Ala His Lys Ile 1000 1005 Gln Ser Ala Asn His Met Asp Asp Asn Asp Thr Glu Leu Asp Thr Pro 1015 1020 Ile Asn Tyr Ser Leu Lys Tyr Ser Asp Glu Gln Leu Asn Ser Gly Arg 1035 1025 1030 Gln Ser Pro Thr Gln Asn Glu Arg Trp Ser Arg Pro Lys His Ile Ile 1045 1050 Asp Ser Glu Met Lys Gln Ser Glu Gln Arg Gln Pro Arg Thr Thr Lys 1060 1065 1070 Thr Thr Tyr Ser Ser Tyr Thr Glu Asn Lys Glu Glu Lys His Lys Lys 1075 1080 1085 Phe Pro Pro His Phe Asn Gln Ser Glu Asn Val Pro Ala Tyr Thr Arg 1090 1095 1100 Ser Arg Gly Ala Asn Asn Gln Val Asp Gln Ser Arg Val Ser Ser Asn 1110 1115 Leu Ser Asn Asn Ser Lys Ala Ser Lys Pro His Cys Gln Val Asp Asp 1125 1130 Tyr Asp Asp Asp Lys Thr Thr Asn Phe Ser Glu Arg Tyr Ser Glu Glu 1145 Glu Gln Gln Glu Asp Glu Thr Glu Arg Gln Asn Lys Tyr Asn Ile Lys 1160 1165 Ala Tyr Ala Ser Glu Glu His His Gly Glu Gln Pro Ile Asp Tyr Ser 1175 1180 Arg Lys Tyr Ser Thr Asp Val Pro Ser Ser Ala Gln Lys Pro Ser Phe 1190 1195 Pro Tyr Ser Asn Asn Ser Ser Lys Gln Lys Pro Lys Lys Glu Gln Val 1205 1210 Ser Ser Asn Ser Asn Thr Pro Thr Pro Ser Pro Asn Ser Asn Arg Gln

122	0	1225	1230	
Asn Gln Leu His 1235	Pro Asn Ser	Ala Gln Ser A 1240	Arg Pro Gly Leu 1245	Asn Arg
Pro Lys Gln Ile 1250	Pro Asn Lys 125		Ile Asn Gln Glu 1260	Thr Ile
Gln Thr Tyr Cys 1265	Val Glu Asp 1270		Cys Phe Ser Arg 1275	Gly Ser 1280
Ser Leu Ser Ser	Leu Ser Ser 1285	Ala Glu Asp (1290	Glu Ile Glu Gly	Arg Glu 1295
Arg Asn Ser Arg 130	_	Ser Asn Asn 5	Thr Leu Gln Ile 1310	
Pro Lys Glu Ile 1315	Ser Ala Val	Ser Lys Asp (Gly Ala Val Asn 1325	Glu Thr
Arg Ser Ser Val 1330	His His Thr 133		Asn Asn Arg Leu 1340	Gln Thr
1345	1350		His Lys Ser Val 1355	1360
	1365	1370		1375
138	0	1385	Pro Leu Met Phe 1390	-
1395		1400	Glu Ser His Ser 1405	
1410	141	5	His Met Ile Ser 1420	
1425	1430	:	Gly Gln Thr Met 1435	1440
	1445	1450		1455
Asp Gly Ser Lys 146		Pro Asp Glu (1465	Glu Arg Gly Lys 1470	
Lys Thr Ala Val 1475	His Ser Ala	Ile Gln Arg V	Val Gln Val Leu 1485	Gln Glu
Ala Asp Thr Leu 1490	Leu His Phe 149		Ser Thr Pro Asp 1500	Gly Phe
1505	1510		Léu Asp Glu Pro 1515	1520
	1525	1530	Pro Val Leu Glu	1535
154	0	1545	Phe Ile Asp Asn 1550	-
1555		1560	Lys Asp Met Leu 1565	
1570	157	5	Cys Ile Ile Ser 1580	
1585	1590		Val Pro Gln Pro 1595	1600
	1605	1610	Pro Ser Gln Leu	1615
162	0	1625	Gln Thr Gln Lys 1630	
1635		1640	Val Tyr Cys Val 1645	
1650	165	5	Leu Ser Asp Leu 1660	
1665	1670		Gln Pro Asn Thr 1675	1680
	1685	1690		1695
Ser Thr Asp Asp	Thr Asp Ala	Ser Lys Pro	Leu Asn Pro Thr	Thr Val

1700	2	1705		1710	
Leu Asp Glu Asp 1715			sp Ile Leu		
His Ser Ala Met 1730	Pro Lys Gly 173		is Lys Pro 1740		
Lys Ile Met Asp 1745	Gln Ile Asn 1750	His Thr Se	er Ala Ala 1755	Thr Ser Ser Gly 1760	o
Asn Ser Arg Ser	Met Gln Glu 1765		ys Asn Lys 770	Pro Thr Ser Pro 1775	
Val Lys Pro Met 1780		Ile Gly Ph 1785	he Lys Glu	Arg Leu Lys Lys 1790	
Asn Thr Glu Leu 1795	Lys Leu Asn	Pro Asn Se	er Glu Asn	Gln Tyr Cys Asp 1805	
Pro Arg Lys Pro	Ser Ser Lys	Lys Pro Se	er Lys Val	Ala Asn Glu Lys	
1810	181	5	1820	ı	
Ile Pro Asn Asn 1825	Glu Glu Arg 1830	Thr Lys G	ly Phe Ala 1835	Phe Asp Ser Pro	n
His His Tyr Thr					•
Asp Ser Leu Ser 1860	Ser Leu Asp	Phe Glu As 1865	sp Asp Asp	Ile Asp Leu Ser 1870	
Lys Glu Lys Ala 1875	Glu Leu Arg	Lys Glu Ly 1880	ys Gly Thr	Lys Asp Thr Asp 1885	
Gln Lys Val Lys 1890	Tyr Lys His 189		rg Ala Ile 1900	——————————————————————————————————————	
Lys Gln Asp Gln 1905	Thr Gly Pro 1910	Lys Ser Le	eu Gly Gly 1915	Arg Asp Gln Pro 1920	o
Lys Ala Leu Val	Gln Lys Pro 1925		he Ser Ser 930	Ala Ala Lys Gly 1935	
Thr Gln Asp Arg	o	1945		1950	
Ile Glu Asn Thr 1955		1960		1965	
Leu Ser Asp Ile 1970	197	5	1980		
Lys Gln Thr Gly 1985	1990		1995	2000	
Thr Ser Gly Tyr	2005	20	010	2015	
Cys Phe Ser Arg	0	2025		2030	
Asp Asp Leu Leu 2035	_	2040		2045	
Lys Pro Ser Lys 2050	205	5	2060	1	
Val Gly Gly Ile 2065	2070		2075	2080	
Asp Ile Gln Ser	Pro Asp Ser 2085		la Phe Ser 090	Pro Asp Ser Glu 2095	
Asn Phe Asp Trp 210	0	2105		2110	
Arg Leu His Gln 2115	Ala Ala Ala	Ala Gly Se 2120	er Leu Ser	Arg Gln Gly Ser 2125	
Ser Asp Ser Asp 2130	213	5	2140		
Ser Pro Phe His 2145	Leu Thr Leu 2150	Asp Lys G	lu Glu Lys 2155	Thr Ile Thr Ser 2160	
Asn Lys Gly Pro	Lys Ile Leu 2165		la Glu Lys 170	Ser Ala Leu Glu 2175	
Asn Lys Lys Thr	Glu Glu Glu	Pro Lys G	ly Ile Lys	Gly Gly Lys Lys	

	2180			2185)				2190	1	
Val Tyr Lys 2195			Gly 2200		Ser	Arg	Ser	Ser 2205		Asp	Phe
Ser Ser His 2210	Cys Lys G	Gln Ser 2215		Gln	Thr	Asn	Met 2220		Ser	Ile	Ser
Arg Gly Arg 2225		le His 2230	Ile	Pro	Gly	Val 2235		Ala	Ser	Ser	Pro 2240
Ser Thr Ser	Pro Val S 2245	Ser Lys	Lys	_	Pro 2250		Phe	Lys	Asn	Val 2255	
Ser Lys Gly	Ser Asn G 2260	Glu Asn		Ser 2265		Ser	Ser		Pro 2270		Gly
Thr Lys Pro 2275	_	Ser Glu	Leu 2280		Tyr	Gly	Ser	Arg 2285		Ser	Ser
Thr Pro Gly 2290	Gly Ser S	Ser Lys 2295		Asn	Ser	Arg	Ser 2300		Ser	Arg	Asp
Ser Ala Ser 2305		Pro Ser 2310	Pro	Gln	Pro	Leu 2315		Arg	Pro	Leu	Gln 2320
Ser Pro Gly	Arg Asn S 2325	Ser Ile	Ser		Gly 2330		Asn	Gly	Ile	Ser 2335	
	2340			2345	<u>,</u>				2350)	
Ser Thr Lys 2355			2360)				2365	•		
Arg Gln Leu 2370		2375	5				2380)			
Thr His Ser 2385	2	2390				2395	•	_			2400
Gln Asn Val	2405	_		_	2410)				2415	5
	2420	_		2425	,	_			2430)	
Ala Leu Val 2435	_		2440)	_			2445	,		
Leu Arg Arg 2450		2455	5				2460)			
Ser Ser Arg 2465	2	2470				2475	,				2480
Leu Ser Pro	2485				2490)				2495	5
	2500			2505	5				2510)	
His Gly Asp 2515			2520)				2525	•		
Ser Pro Ser 2530		2535	5				2540)			
His Ser Lys 2545	2	2550				2555	•			-	2560
Thr Gly Ser	2565				2570)				2575	5
	2580		_	2585	5		_		2590)	<u></u>
Pro Arg Ser 2595	_		2600)		_		2605	,		
Lys Glu Ser 2610		2615	5				2620)			
Ile Ala Glu 2625	2	2630				2635	·)			_	2640
Met Ala Pro	2645				2650)				2655	5
Asp Cys Pro	Ile Asn A 2660	Asn Pro	Arg	Ser 2665		Arg	Ser	Pro	Thr 2670		Asn

Ser	Pro	Pro 2675		Ile	Asp	Asn	Val 2680		Asp	Gln	Gly	Gln 2685		Glu	Glu	
Ala	Ala 2690		Asp	Cys	His	Thr 2699		His	Asn	Ser	Gly 2700		Gly	Asn	Val	
Pro 2705		Leu	Glu	Asn	Arg 2710		Lys	Ser	Phe	Ile 2715		Val	Asp	Gly	Leu 2720	
Asp	Thr	Lys	Gly	Thr 2725		Pro	Lys	Ser	Leu 2730		Asn	Asn	Gln	Gln 2735		
Thr	Asn	Glu	Asn 2740		Val	Ala	Glu	Arg 2745		Ala	Phe	Ser	Ser 2750	Ser	Ser	
Ser	Ser	Lys 2755		Ser	Ser	Pro	Ser 2760	_	Thr	Val	Ala	Ala 2765	_	Val	Thr	
Pro	Phe 2770		Tyr	Asn	Pro	Ser 2775		Arg	Lys	Ser	Asn 2780	-	Glu	Asn	Ser	
Thr 2785		Arg	Pro	Ser	Gln 2790		Pro	Thr	Pro	Val 2795		Asn	Ser	Thr	Lys 2800	
Lys	Arg	Asp	Ser	Lys 2805		Glu	Thr	Thr	Asp 2810		Ser	Gly	Ser	Gln 2815		
Pro	Lys	Arg	His 2820		Gly	Ser	Tyr	Leu 2825		Thr	Ser	Val				
<210																
	.> 30 ?> DN															
			icial	l Sec	queno	ce										
<220)>															
<223	3> Ar	citif	fical	lly s	syntl	nesi	zed p	prime	er se	equer	nce					
<400)> 2															
cgac	gcgt	aa t	gcat	tttt	ct co	caga	ctct	3								30
<210)> 3															
	> 39															
	?> Dî ?> Aî		icial	l Sed	quenc	ce										
					•											
<220 <223		ritif	fical	lly s	syntl	nesi	zed p	prime	er se	equer	nce					
<400												•				
ggaa	ittc	gga t	cct	cacao	cc aç	gataa	agaad	c caq	gagt	gcc						39
<210)> 4															
	.> 33															
	?> Dì		1) C												
<213	5> A1	CLIII	ıcıaı	ı Sed	queno	ce										
<220 <223		ritif	fical	lly s	synth	nesi:	zed p	orime	er se	equer	nce					
<400)> 1															
		at q	ggct	gctgo	et to	gtai	gate	c agt	Ξ.							33
<210)> 5															
	.> 29												•			
	?> Dî ?> Ar		icia	l Sec	quenc	ce										
~					· · ·	-										

<220>	
<223> Aritifically synthesized primer sequence	
<400> 5 cgacgcgtac ctgctgttct ttccctgtc	29
<210> 6 <211> 28 <212> DNA <213> Artificial Sequence	
<220> <223> Aritifically synthesized primer sequence	
<400> 6 ctagctagca tggctgctgc ttcgtatg	28
<210> 7 <211> 27 <212> DNA <213> Artificial Sequence	
<220> <223> Aritifically synthesized primer sequence	
<400> 7 cctgtcccaa gtaggtcacg atcgatc	27
<210> 8 <211> 27 <212> DNA <213> Artificial Sequence	
<220> <223> Aritifically synthesized primer sequence	
<400> 8 ctagctagcc tcggcaacta ccattcg	27
<210> 9 <211> 19 <212> DNA <213> Artificial Sequence	
<220> <223> Aritifically synthesized primer sequence	
<400> 9 attagagete actetagae	19